



033488-001.ST25

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<110> Weber, Bernard H.F.
Stoehr, Heidi

<120> Novel retina-specific human proteins C7orf9, C12orf7, MPP4 and F379

<130> 033488-001

<140> US 09/995,793

<141> 2001-11-29

<150> 60/253,751

<151> 2000-11-29

<160> 71

<170> PatentIn version 3.1

<210> 1

<211> 2435

<212> DNA

<213> Homo sapiens

<220>

<221> misc feature

<223> artificial sequence, Translation start at 209; stop at 2435

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<220>
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<210> 4
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<212> DNA
 <213> Homo sapiens

<220>
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 <223> genomic DNA, Exon from 165 to 286

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 ctgtaggcca ggaagtccaa aatccagt 448

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 <211> 448
 <212> DNA
 <213> Homo sapiens

<220>
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 <223> genomic DNA, Exon from 206 to 283

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<210> 6
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 <211> 448
 <212> DNA
 <213> Homo sapiens

<220>
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 <223> genomic DNA, Exon from 133 to 264

<400> 7
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 <213> Homo sapiens

<220>
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 <223> genomic DNA, Exon from 166 to 247

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<210> 9
 <211> 448
 <212> DNA
 <213> Homo sapiens

<220>
 <221> misc_feature
 <223> genomic DNA, Exon from 162 to 247

<400> 9
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448

<210> 10
 <211> 384
 <212> DNA
 <213> Homo sapiens

<220>
 <221> misc_feature
 <223> genomic DNA, Exon from 158 to 229

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 aatactgatg gcccgagaag caaa 384

<210> 11
 <211> 448
 <212> DNA
 <213> Homo sapiens

<220>
 <221> misc_feature
 <223> genomic DNA, Exon from 138 to 334

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 ataaagaggg acaagcagat gaatgaac 448

<210> 12
 <211> 320
 <212> DNA
 <213> Homo sapiens

<220>
 <221> misc_feature
 <223> genomic DNA, Exon from 152 to 216

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 cgtaccagcc tcacacctgc ctcaagtcaa ccctatgtga gtattgcaac tgcccagacg 240
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<210> 13
 <211> 320
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 <213> Homo sapiens

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 aagaaatagt atttaggaaa aaactcttat ctccaaagtc ttttagaaat ttctttagt 240
 ttaaagaatt cactttaatt cagttcagct atttattaag ctcttcctat atacctagta 300
 gtgtgatagt cattattaag 320

<210> 14
 <211> 384
 <212> DNA
 <213> Homo sapiens

<220>
 <221> misc_feature
 <223> genomic DNA, Exon from 179 to 217

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 <212> DNA
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<210> 16
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<210> 17
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<210> 18
 <211> 512
 <212> DNA
 <213> Homo sapiens

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<210> 19
 <211> 384
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 <213> Homo sapiens

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 cacatctgat gatttctgtg tgtgactttt tgtgttttagg accctctggt gttggagtaa 180
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<210> 20
 <211> 448
 <212> DNA
 <213> Homo sapiens

<220>
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 <223> genomic DNA, Exon from 200 to 293

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 agatatcttt attcaaattgc atattggtta tcaaagaatt ctgaagacac tgaaaccttt 180
 cattcccttt ttctgataga cactactcgt actaaaaaga gttacgaaat gaatgggcgt 240
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<210> 21
 <211> 448
 <212> DNA
 <213> Homo sapiens

<220>
 <221> misc_feature
 <223> genomic DNA, Exon from 133 to 241

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<210> 22
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<213> Homo sapiens

<220>

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<223> genomic DNA, Exon from 164 to 298

<400> 22

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<210> 23

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<212> DNA

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<400> 23

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acagcataac aaactgtatt ttttccattt gtccaattaa gtctgtacta tccatatttt      180
tctattttct ctaaaggatg aagacctaca agagatggaa aatttagccc aaagaatgga      240
aactcagttt ggccaatttt ttgatcatgt gattgtgaat gacagcttgc acgatgcatg      300
tgcccagttg ttgtctgcca tacagaaggc tcaggaggag cctcagtggg taccagcaac      360
atggatttcc tcagatactg agtctcaatg agacttcttg tttaatgctg gagttttaac      420
actgtaccct tgatacagcg atccatagtt gcaatctaaa acaacagtat ttgacccatt      480
ttaatgtgta caactttaaa agtgcagcaa tttattaatt aatcttattt gaaaaaaatt      540
tttattgtat ggttatgtgg ttacctatgt taacttaatt ttttttcctt tacctcatat      600
gcagctgtgg tagaaatatg aataatgtta agtcactgag tatgagaacc tttcgcagat      660
ttcacatgat ctttttaaga tttaaataaa gagctttcct aaat                                     704

```

<210> 24

<211> 637

<212> PRT

<213> Homo sapiens

<400> 24

```

Met Ile Gln Ser Asp Lys Gly Ala Asp Pro Pro Asp Lys Lys Asp Met
1           5           10           15
Lys Leu Ser Thr Ala Thr Asn Pro Gln Asn Gly Leu Ser Gln Ile Leu
20           25           30
Arg Leu Val Leu Gln Glu Leu Ser Leu Phe Tyr Ser Arg Asp Val Asn
35           40           45
Gly Val Cys Leu Leu Tyr Asp Leu Leu His Ser Pro Trp Leu Gln Ala
50           55           60
Leu Leu Lys Ile Tyr Asp Cys Leu Gln Glu Phe Lys Glu Lys Lys Leu

```

65					70					75					80
Val	Pro	Ala	Thr	Pro	His	Ala	Gln	Val	Leu	Ser	Tyr	Glu	Val	Val	Glu
				85					90					95	
Leu	Leu	Arg	Glu	Thr	Pro	Thr	Ser	Pro	Glu	Ile	Gln	Glu	Leu	Arg	Gln
			100					105					110		
Met	Leu	Gln	Ala	Pro	His	Phe	Lys	Ala	Leu	Leu	Ser	Ala	His	Asp	Thr
		115					120					125			
Ile	Ala	Gln	Lys	Asp	Phe	Glu	Pro	Leu	Leu	Pro	Pro	Leu	Pro	Asp	Asn
	130						135				140				
Ile	Pro	Glu	Ser	Glu	Glu	Ala	Met	Arg	Ile	Val	Cys	Leu	Val	Lys	Asn
145					150					155				160	
Gln	Gln	Pro	Leu	Gly	Ala	Thr	Ile	Lys	Arg	His	Glu	Met	Thr	Gly	Asp
				165					170					175	
Ile	Leu	Val	Ala	Arg	Ile	Ile	His	Gly	Gly	Leu	Ala	Glu	Arg	Ser	Gly
		180						185					190		
Leu	Leu	Tyr	Ala	Gly	Asp	Lys	Leu	Val	Glu	Val	Asn	Gly	Val	Ser	Val
	195					200					205				
Glu	Gly	Leu	Asp	Pro	Glu	Gln	Val	Ile	His	Ile	Leu	Ala	Met	Ser	Arg
	210					215				220					
Gly	Thr	Ile	Met	Phe	Lys	Val	Val	Pro	Val	Ser	Asp	Pro	Pro	Val	Asn
225					230					235				240	
Ser	Gln	Gln	Met	Val	Tyr	Val	Arg	Ala	Met	Thr	Glu	Tyr	Trp	Pro	Gln
			245						250					255	
Glu	Asp	Pro	Asp	Ile	Pro	Cys	Met	Asp	Ala	Gly	Leu	Pro	Phe	Gln	Lys
		260						265					270		
Gly	Asp	Ile	Leu	Gln	Ile	Val	Asp	Gln	Asn	Asp	Ala	Leu	Trp	Trp	Gln
	275						280					285			
Ala	Arg	Lys	Ile	Ser	Asp	Pro	Ala	Thr	Cys	Ala	Gly	Leu	Val	Pro	Ser
	290					295					300				
Asn	His	Leu	Leu	Lys	Arg	Lys	Gln	Arg	Glu	Phe	Trp	Trp	Ser	Gln	Pro
305					310					315				320	
Tyr	Gln	Pro	His	Thr	Cys	Leu	Lys	Ser	Thr	Leu	Ser	Ile	Ser	Met	Glu
			325						330					335	
Glu	Glu	Asp	Asp	Met	Lys	Ile	Asp	Glu	Lys	Cys	Val	Glu	Ala	Asp	Glu
		340						345					350		
Glu	Thr	Phe	Glu	Ser	Glu	Glu	Leu	Ser	Glu	Asp	Lys	Glu	Glu	Phe	Val
	355						360					365			
Gly	Tyr	Gly	Gln	Lys	Phe	Phe	Ile	Ala	Gly	Phe	Arg	Arg	Ser	Met	Arg
	370					375				380					
Leu	Cys	Arg	Arg	Lys	Ser	His	Leu	Ser	Pro	Leu	His	Ala	Ser	Val	Cys
385					390					395				400	
Cys	Thr	Gly	Ser	Cys	Tyr	Ser	Ala	Val	Gly	Ala	Pro	Tyr	Glu	Glu	Val
			405						410					415	
Val	Arg	Tyr	Gln	Arg	Arg	Pro	Ser	Asp	Lys	Tyr	Arg	Leu	Ile	Val	Leu
		420						425					430		
Met	Gly	Pro	Ser	Gly	Val	Gly	Val	Asn	Glu	Leu	Arg	Arg	Gln	Leu	Ile
	435						440					445			
Glu	Phe	Asn	Pro	Ser	His	Phe	Gln	Ser	Ala	Val	Pro	His	Thr	Thr	Arg
	450					455					460				
Thr	Lys	Lys	Ser	Tyr	Glu	Met	Asn	Gly	Arg	Glu	Tyr	His	Tyr	Val	Ser
465					470					475				480	
Lys	Glu	Thr	Phe	Glu	Asn	Leu	Ile	Tyr	Ser	His	Arg	Met	Leu	Glu	Tyr
			485						490					495	
Gly	Glu	Tyr	Lys	Gly	His	Leu	Tyr	Gly	Thr	Ser	Val	Asp	Ala	Val	Gln
		500						505				510			
Thr	Val	Leu	Val	Glu	Gly	Lys	Ile	Cys	Val	Met	Asp	Leu	Glu	Pro	Gln
	515						520					525			

```

Asp Ile Gln Gly Val Arg Thr His Glu Leu Lys Pro Tyr Val Ile Phe
530                               535                               540
Ile Lys Pro Ser Asn Met Arg Cys Met Lys Gln Ser Arg Lys Asn Ala
545                               550                               555                               560
Lys Val Ile Thr Asp Tyr Tyr Val Asp Met Lys Phe Lys Asp Glu Asp
565                               570                               575
Leu Gln Glu Met Glu Asn Leu Ala Gln Arg Met Glu Thr Gln Phe Gly
580                               585                               590
Gln Phe Phe Asp His Val Ile Val Asn Asp Ser Leu His Asp Ala Cys
595                               600                               605
Ala Gln Leu Leu Ser Ala Ile Gln Lys Ala Gln Glu Glu Pro Gln Trp
610                               615                               620
Val Pro Ala Thr Trp Ile Ser Ser Asp Thr Glu Ser Gln
625                               630                               635

```

```

<210> 25
<211> 1190
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<223> artificial sequence, Translation start at 48, stop at 638

```

```

<400> 25
ataaacattg ggctgcacat agagacttaa ttttagattt agacaaaatg gaaattatatt 60
catcaaaact attcatttta ttgacttttag ccacttcaag cttgttaaca tcaaacattt 120
tttgtgcaga tgaattagtg atstccaatc ttcacagcaa agaaaattat gacaaatatt 180
ctgagcctag aggataccca aaaggggaaa gaagcctcaa ttttgaggaa ttaaaagatt 240
ggggaccaa aaatgttatt aagatgagta cacctgcagt caataaaatg ccacactcct 300
tcgccaactt gccattgaga ttggggagga acgttcaaga agaaagaagt gctggagcaa 360
cagccaacct gcctctgaga tctggaagaa atatggaggt gagcctcgtg agacgtgttc 420
ctaacctgcc ccaaagggtt gggagaacaa caacagccaa aagtgtctgc aggatgctga 480
gtgattttgtg tcaaggatcc atgcattcac catgtgccaa tgacttattt tactccatga 540
cctgccagca ccaagaaatc cagaatcccg atcaaaaaca gtcaaggaga ctgctattca 600
agaaaaataga tgatgcagaa ttgaaacaag aaaaataaga aacctggagc ctgtccctaa 660
agctgtggcc tgtaattctac aaatggctct atagcgaaga ccacacggaa gagtagctac 720
atacacttca tcagctatgg atcatcaacg gcaatttttc cttgtcagta cagctataat 780
agtatcttga aagttgtaaa aaaattaaag catatttggt acgtaaagtt aaaatgattt 840
ttgtctgaat aaaaaaaaaa cattgcaaat gctttagaaa tctctgataa tggagagaga 900
gacagaggac cctcctcact accctatata aaaatcattg gcacagttac acttaataaa 960
aaaaattaaa cagaagagca ccctgaaaaa cattatgatg gaaattaaat agtatgccag 1020
aataacatgg ttgacaaata agtgaacaag gattaaaaat cacttacaaa cgtgtttctg 1080
tacacccttt ctatcgtgtc aaatgttaat gaatctgtga tcaattgaaa tgtaaattgc 1140
tgtgtaaaac tacaaaataa aaactcttag actttaggga gaaaagaaaa 1190

```

```

<210> 26
<211> 256
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<223> genomic DNA, Exon from 1 to 185

```

```

<400> 26
ataaacattg ggctgcacat agagacttaa ttttagattt agacaaaatg gaaattattt 60
catcaaaaact attcatttta ttgacttttag ccacttcaag cttgttaaca tcaaacattt 120
tttgtgcaga tgaattagtg atstccaatc ttcacagcaa agaaaattat gacaaatatt 180
ctgaggtaag ttttttaaat ctctctaag tgagtagcat taattacata atattaatcc 240
taagtctaag gatttt 256

```

```

<210> 27
<211> 512
<212> DNA
<213> Homo sapiens

```

```

<220>
<221> misc_feature
<223> genomic DNA, Exon from 62 to 462

```

```

<400> 27
gggtttaaat ctgttgctta taacaacagt atgttattgt aatgggcatt tctaattata 60
gcctagagga tacccaaaag gggaaagaag cctcaatttt gaggaattaa aagattgggg 120
accaaaaaat gttattaaga tgagtacacc tgcagtcaat aaaatgccac actccttcgc 180
caacttgcca ttgagatttg ggaggaacgt tcaagaagaa agaagtgtg gagcaacagc 240
caacctgcct ctgagatctg gaagaaatat ggaggtgagc ctctgagagc gtgttcctaa 300
cctgccccaa aggtttggga gaacaacaac agccaaaagt gtctgcagga tgctgagtga 360
tttgtgtcaa ggatccatgc attcaccatg tgccaatgac ttattttact ccatgacctg 420
ccagcaccaa gaaatccaga atcccgatca aaaacagtca aggtaaatac ctggaaacca 480
gtcaaaagtgc atgggcagtt atatataggt gg 512

```

```

<210> 28
<211> 768
<212> DNA
<213> Homo sapiens

```

```

<220>
<221> misc_feature
<223> genomic DNA, Exon from 115 to 718

```

```

<400> 28
acacaattca actcaagtat aattaggcag ttaggactat ggcttgtatt tgtatacaca 60
cttgcattgt gttgttctga tgggtgacaa cattttatatac tgcttacatt ttaggagact 120
gctattcaag aaaatagatg atgcagaatt gaaacaagaa aaataagaaa cctggagcct 180
gtccctaaag ctgtggcctg taatctacaa atggctctat agcgaagacc acacggaaga 240
gtagctacat acacttcac agctatggat catcaacggc aatttttctt tgtcagtaca 300
gctataatag tatcttgaaa gttgtaaaaa aattaaagca tatttggtac gtaaagttaa 360
aatgattttt gtctgaataa aaaaaaagca ttgcaaatgc tttagaaatc tctgataatg 420
gagagagaga cagaggacc tcctcactac cctatataaa aatcattggc acagttacac 480
ttaataaaaa aaattaaaca gaagagcacc ctgaaaaaca ttatgatgga aattaaatag 540
tatgccagaa taacatgggt gacaaaataag tgaacaagga ttaaaaatca cttacaaacg 600
tgtttctgta caccctttct atcgtgtcaa atgttaatga atctgtgatc aattgaaatg 660
taaattgtctg tgtaaaacta caaaataaaa actcttagac tttagggaga aaagaaaaag 720
gcaactatga gttacctctt ttagtgtctc ctctatctac atccagaa 768

```

```

<210> 29
<211> 196
<212> PRT

```

<213> Homo sapiens

<400> 29

```

Met Glu Ile Ile Ser Ser Lys Leu Phe Ile Leu Leu Thr Leu Ala Thr
1           5           10          15
Ser Ser Leu Leu Thr Ser Asn Ile Phe Cys Ala Asp Glu Leu Val Ile
          20          25          30
Ser Asn Leu His Ser Lys Glu Asn Tyr Asp Lys Tyr Ser Glu Pro Arg
          35          40          45
Gly Tyr Pro Lys Gly Glu Arg Ser Leu Asn Phe Glu Glu Leu Lys Asp
          50          55          60
Trp Gly Pro Lys Asn Val Ile Lys Met Ser Thr Pro Ala Val Asn Lys
65          70          75          80
Met Pro His Ser Phe Ala Asn Leu Pro Leu Arg Phe Gly Arg Asn Val
          85          90          95
Gln Glu Glu Arg Ser Ala Gly Ala Thr Ala Asn Leu Pro Leu Arg Ser
          100         105         110
Gly Arg Asn Met Glu Val Ser Leu Val Arg Arg Val Pro Asn Leu Pro
          115         120         125
Gln Arg Phe Gly Arg Thr Thr Thr Ala Lys Ser Val Cys Arg Met Leu
          130         135         140
Ser Asp Leu Cys Gln Gly Ser Met His Ser Pro Cys Ala Asn Asp Leu
145         150         155         160
Phe Tyr Ser Met Thr Cys Gln His Gln Glu Ile Gln Asn Pro Asp Gln
          165         170         175
Lys Gln Ser Arg Arg Leu Leu Phe Lys Lys Ile Asp Asp Ala Glu Leu
          180         185         190
Lys Gln Glu Lys
          195

```

<210> 30

<211> 1188

<212> DNA

<213> Homo sapiens

<220>

<221> misc feature

<223> artificial sequence, Translation start at 347, stop at 604

<400> 30

```

acacacaacg gggtttcggg gctgtggacc ctgtgccagg aaaggaaggg cgcagctcct      60
gcaatgcgga gcagccaggg cagtgggcac caggctttag cctccctttc tcaccctaca      120
gagggcaggc ccttcagctc cattctcctc caaggctgca gagggggcag gaattggggg      180
tgacaggaga gctgtaaggt ctccagtggg tcattctggg cccagagatg ggtgctgaag      240
ctcccacgcc tgcctgtgaa aatggagtcc tctctcacct gggagagcca ggtgctgccc      300
cgagaaggat gcatttatgg cttcrtgaag tctttcctga ccccgatgc tgctgactat      360
agagacaaag tctcactatg ttgctcaggc tggctcttgaa ctctggcct caagcgatcc      420
tcccacctya gcctcccaa gwgttgggat tatagacatg agccactgca cctggccgac      480
cttgggcaag ttcttaaacc cttcaaagcc tcatTTTTct ccaatcayaa aagggaaaga      540
tggtaatatt ttccccwcca aattcttgtc ggatgccctc acagaattga gattatgtac      600
gtaaaacacc aggtgcctaa cccggcacag agcaggaggg ctaagcgtga catccagcac      660
gtggtcagtg gaatccagta ttcctaccca cctctctagt ctcccctcca cccctctccc      720
tttcagaggc accaagctgc ttgtggtcct gtctattccc actccctgcc tgactgaaca      780
ttttctccac ctctgatca tcagcagcag aaactggctg ctcttcctcc tgggtagaca      840
gccagactgt atttcccagc tgcccctgca gtgagatgtg gccatcggag ccagcattgg      900

```

```

ccaatggact ctgcatggga gtgacgcatg cwgcctccag gcttgtccct aaaacctccc 960
acgtgtcctc sgcctgctct tcccacytcc aaggagcacg gcaattgtgg aagacccaga 1020
ttagtgatgg cagaaccata gatgggagga acctgggtcc ctgacttaaa gtatcatgga 1080
tttgatgtt cccttagtga gaaataaact tccattgtgt ttaagccttt atttgtttat 1140
agttggttac agcaactgcc ttcttttaat taaaacactc ctgctgct 1188

```

```

<210> 31
<211> 85
<212> PRT
<213> Homo sapiens

```

```
<400> 31
```

```

Met Leu Leu Thr Ile Glu Thr Lys Ser His Tyr Val Ala Gln Ala Gly
1          5          10          15
Leu Glu Leu Leu Ala Ser Ser Asp Pro Pro Thr Ser Ala Ser Gln Ser
20          25          30
Val Gly Ile Ile Asp Met Ser His Cys Thr Trp Pro Thr Leu Gly Lys
35          40          45
Phe Leu Asn Pro Ser Lys Pro His Phe Ser Pro Ile Thr Lys Gly Lys
50          55          60
Asp Gly Asn Ile Phe Pro Thr Lys Phe Leu Ser Asp Ala Leu Thr Glu
65          70          75          80
Leu Arg Leu Cys Thr
85

```

```

<210> 32
<211> 560
<212> DNA
<213> Homo sapiens

```

```

<220>
<221> misc_feature
<223> genomic DNA, Exon from 101 to 460

```

```

<400> 32
tatatgggaa tgagccagct gcaccgctgc tgacagtggc tgggataatc ctccctgagc 60
tgttccaagg attagtcctg ctgccctgtg cccagctccc acacaacggg gtttcggggc 120
tgtggaccct gtgccaggaa aggaagggcg cagctcctgc aatgcggagc agccagggca 180
gtgggcacca ggcttttagcc tccctttctc accctacaga gggcaggccc ttcagctcca 240
ttctcctcca aggctgcaga gggggcagga attgggggtg acaggagagc tgtaaggtct 300
ccagtgggtc attctgggcc gagagatggg tgctgaagct cccacgcctg cctgtgaaaa 360
tggagtcctc tctcacctgg gagagccagg tgctgccccg agaaggatgc atttatggct 420
tcatgaagtc tttcctgacc cccgatgctg ctgactatag gtaagtctga gcaaactctg 480
gggagcctca tcttgcatg agaaagagat ggcttcttct aagccactg gccgtgatcc 540
caggattata acacattctg
560

```

```

<210> 33
<211> 405
<212> DNA
<213> Homo sapiens

```

```

<220>
<221> misc_feature

```

<223> genomic DNA, Exon from 101 to 305

<400> 33

catgagaggt	agtataatat	agaggatatg	tgtgcttact	aagaggctgc	ctgtctgacc	60
ttggacaagt	tctttttatt	tattttattta	ttttttatag	agacaaagtc	tcactatggt	120
gctcaggctg	gtcttgaact	cctggcctca	agcgatcctc	ccaccttagc	ctcccaaaga	180
gttgggatta	tagacatgag	ccactgcacc	tggccgacct	tgggcaagtt	cttaaaccct	240
tcaaagcctc	atttttctcc	aatcataaaa	gggaaagatg	gtaatatatt	cccctccaaa	300
ttcttgtaag	tattaaacat	tgtatatgta	ttttgaacac	gattaagctc	taaacacttg	360
ttaggaagca	ggagtagcat	ttgaaacaaa	cagctctttt	cccac		405

<210> 34

<211> 821

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<223> genomic DNA, Exon from 101 to 721

<400> 34

aagtattaaa	cattgtatat	gtattttgaa	cacgattaag	ctctaaacac	ttgttaggaa	60
gcaggagtag	catttgaaac	aaacagctct	tttcccacag	gtcggatgcc	ctcacagaat	120
tgagattatg	tacgtaaaac	accagggtgcc	taaccoggca	cagagcagga	gggctaagcg	180
tgacatccag	cacgtggtca	gtggaatcca	gtattcctac	ccacctctct	agtctccctt	240
ccacccctct	ccctttcaga	ggcaccaagc	tgcttgtggt	cttgtctatt	cccactccct	300
gcctgactga	acattttctc	cacctcctga	tcatcagcag	cagaaactgg	ctgctcttcc	360
tcctgggtag	acagccagac	tgtatttccc	agctgcccct	gcagtgagat	gtggccatcg	420
gagccagcat	tggccaatgg	actctgcatg	ggagtgacgc	atgctgcctc	caggcttgtc	480
cctaaaacct	cccacgtgtc	ctccgcctgc	tcttcccact	tccaaggagc	acggcaattg	540
tggaagaccc	agattagtga	tggcagaacc	atagatggga	ggaacctggg	tccctgactt	600
aaagtatcat	ggattttggat	gttcccttag	tgagaaataa	acttccattg	tgtttaagcc	660
tttatttggt	tatagttggt	tacagcaact	gcctttcttt	aattaaaaca	ctcctgctgc	720
ttcatgttgc	tggaatgctt	gtaaccctgc	cctgcttcac	cagggtaact	cctacttggc	780
ctttaagttt	atctctgctg	tcacaccgtc	cagaaagcct	t		821

<210> 35

<211> 1514

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<223> artificial sequence, Translation start at 155, stop at 1192

<400> 35

gaaagtccag	ccatctgtta	cctgcgttgc	ttcctggggr	gggatagtcc	acctggaggc	60
attcggagac	ccagtgattg	tgtccgygg	agcctgggct	gtgccccgcg	ttgactgcct	120
catagatacc	ctacgaaccc	caaattgccag	ctgcatgaga	aaagggactc	accttctggt	180
tccctgcctg	gaagaggaag	agctggcatt	gcacaggaga	cggctggaca	tgtctgaggc	240
actgccctgc	ccgggcaagg	agacccccac	cccaggctgc	aggctggggg	ccctgtattg	300
ggcctgtgtc	cacaatgac	ccaccagct	ccaagccata	ctggatgggtg	gggtctcccc	360
agaggaggcc	accaggtgg	acagcaatgg	gaggacaggc	ctcatgggtcg	catgctacca	420
cggcttccag	agtgttggtg	ccctgctcag	ccactgtcct	ttccttgatg	tgaaccagca	480
ggacaaagga	ggggacacgg	ccctcatgtt	ggctgcccac	gcaggccacg	tgccctctagt	540

gagtctcctg	ctcaactact	atgtgggcct	ggacctggaa	cgccgggacc	agcgggggct	600
cacggcggtta	atgaaggctg	ccatgcggaa	ccgctgtgct	gacctgacag	cagtggaccc	660
tgttcggggc	aagacggccc	tggaaatgggc	agtgtgacc	gacagcttcg	acaccgtgtg	720
gaggattcgg	cagctgctga	ggcggcccca	agtggagcag	cttagccagc	actacaagcc	780
cgagtggccg	gccttgccg	ggctcgtggc	ccaggcccag	gcccaggccc	aggttgcccc	840
ttcactccta	gaacggctgc	aggctacctt	gagcctcccc	tttgccccgt	ctcctcagga	900
gggggggtgtt	ctggaccacc	ttgtgactgc	cacaaccagc	ctggccagtc	ccttcgtcac	960
cactgcctgc	cacactctgt	gccctgacca	tccaccttcg	ctgggcaccc	gaagcaagtc	1020
cgtgccagag	ctgttagtgc	cagccgaagc	ccagtccttc	aggacaccaa	agtctggccc	1080
ttcctctctg	gcgataccag	gagctcagga	tagagaagag	gaaacaggag	gaggaggcca	1140
gaatggcaca	gaagtagggg	aagatgggat	aggacaggct	gggaacaggt	aatcaggccc	1200
ctcccagggc	ttctttcccc	tctggagtgc	ctccggcctc	cccatccacc	tctgcctaag	1260
taaatctgct	ctcaacctat	atatatacaa	ggtcattcat	tctagcattg	tttgaagag	1320
tgaagagtg	gaaacacccg	aagtgtccat	cagtaaggga	caggctagat	tgattacgga	1380
tgtaattgct	gtccatccat	acagagcata	ctctacagtg	tattctaaaa	taagactaag	1440
gaagctgttt	atattctgat	atgaaactac	catcaagatg	tataaagtaa	aaataactaa	1500
ggagtgaac	agtg					1514

<210> 36

<211> 1544

<212> DNA

<213> Homo sapiens

<220>

<221> misc feature

<223> artificial sequence, Translation start at 155, stop at 1222

<400> 36

gaaagtccag	ccatctgtta	cctgcgttgc	ttcctggggr	gggatagtcc	acctggaggc	60
attcggagac	ccagtgattg	tgctccgygg	agcctgggct	gtgcccgcg	ttgactgcct	120
catagatacc	ctacgaaccc	caaatgccag	ctgcatgaga	aaagggactc	accttctggt	180
tccttgccctg	gaagaggaag	agctggcatt	gcacaggaga	cggctggaca	tgtctgaggc	240
actgccctgc	ccgggcaagg	agacccccac	cccaggctgc	aggctggggg	ccctgtattg	300
ggcctgtgtc	cacaatgatc	ccaccagct	ccaagccata	ctggatggtg	gggtctcccc	360
agaggaggcc	accaggtgg	acagcaatgg	gaggacaggc	ctcatggtcg	catgctacca	420
cggcttcag	agtgttgtg	ccctgctcag	ccactgtcct	ttccttgatg	tgaaccagca	480
ggacaaaagga	ggggacacgg	ccctcatgtt	ggctgcccac	gcaggccacg	tgctcttagt	540
gagtctcctg	ctcaactact	atgtgggcct	ggacctggaa	cgccgggacc	agcgggggct	600
cacggcggtta	atgaaggctg	ccatgcggaa	ccgctgtgag	tgctggcca	ccctcctcat	660
ggcaggtgct	gacctgacag	cagtggaccc	tgctcggggc	aagacggccc	tggaaatgggc	720
agtgtgacc	gacagcttcg	acaccgtgtg	gaggattcgg	cagctgctga	ggcggcccca	780
agtggagcag	cttagccagc	actacaagcc	cgagtggccg	gccttgccg	ggctcgtggc	840
ccaggcccag	gcccaggccc	aggttgcccc	ttcactccta	gaacggctgc	aggctacctt	900
gagcctcccc	tttgccccgt	ctcctcagga	gggggggtgtt	ctggaccacc	ttgtgactgc	960
cacaaccagc	ctggccagtc	ccttcgtcac	cactgcctgc	cacactctgt	gccctgacca	1020
tccaccttcg	ctgggcaccc	gaagcaagtc	cgtgccagag	ctgttagtgc	cagccgaagc	1080
ccagtccttc	aggacaccaa	agtctggccc	ttcctctctg	gcgataccag	gagctcagga	1140
tagagaagag	gaaacaggag	gaggaggcca	gaatggcaca	gaagtagggg	aagatgggat	1200
aggacaggct	gggaacaggt	aatcaggccc	ctcccagggc	ttctttcccc	tctggagtgc	1260
ctccggcctc	cccatccacc	tctgcctaag	taaatctgct	ctcaacctat	atatatacaa	1320
ggtcattcat	tctagcattg	tttgaagag	tgaagagtg	gaaacacccg	aagtgtccat	1380
cagtaaggga	caggctagat	tgattacgga	tgtaattgct	gtccatccat	acagagcata	1440
ctctacagtg	tattctaaaa	taagactaag	gaagctgttt	atattctgat	atgaaactac	1500
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<210> 37
 <211> 345
 <212> PRT
 <213> Homo sapiens

<400> 37

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Met Arg Lys Gly Thr His Leu Leu Val Pro Cys Leu Glu Glu Glu Glu
1      5      10      15
Leu Ala Leu His Arg Arg Arg Leu Asp Met Ser Glu Ala Leu Pro Cys
20      25      30
Pro Gly Lys Glu Thr Pro Thr Pro Gly Cys Arg Leu Gly Ala Leu Tyr
35      40      45
Trp Ala Cys Val His Asn Asp Pro Thr Gln Leu Gln Ala Ile Leu Asp
50      55      60
Gly Gly Val Ser Pro Glu Glu Ala Thr Gln Val Asp Ser Asn Gly Arg
65      70      75      80
Thr Gly Leu Met Val Ala Cys Tyr His Gly Phe Gln Ser Val Val Ala
85      90      95
Leu Leu Ser His Cys Pro Phe Leu Asp Val Asn Gln Gln Asp Lys Gly
100     105     110
Gly Asp Thr Ala Leu Met Leu Ala Ala Gln Ala Gly His Val Pro Leu
115     120     125
Val Ser Leu Leu Leu Asn Tyr Tyr Val Gly Leu Asp Leu Glu Arg Arg
130     135     140
Asp Gln Arg Gly Leu Thr Ala Leu Met Lys Ala Ala Met Arg Asn Arg
145     150     155     160
Cys Ala Asp Leu Thr Ala Val Asp Pro Val Arg Gly Lys Thr Ala Leu
165     170     175
Glu Trp Ala Val Leu Thr Asp Ser Phe Asp Thr Val Trp Arg Ile Arg
180     185     190
Gln Leu Leu Arg Arg Pro Gln Val Glu Gln Leu Ser Gln His Tyr Lys
195     200     205
Pro Glu Trp Pro Ala Leu Ser Gly Leu Val Ala Gln Ala Gln Ala Gln
210     215     220
Ala Gln Val Ala Pro Ser Leu Leu Glu Arg Leu Gln Ala Thr Leu Ser
225     230     235     240
Leu Pro Phe Ala Pro Ser Pro Gln Glu Gly Gly Val Leu Asp His Leu
245     250     255
Val Thr Ala Thr Thr Ser Leu Ala Ser Pro Phe Val Thr Thr Ala Cys
260     265     270
His Thr Leu Cys Pro Asp His Pro Pro Ser Leu Gly Thr Arg Ser Lys
275     280     285
Ser Val Pro Glu Leu Leu Val Pro Ala Glu Ala Gln Ser Phe Arg Thr
290     295     300
Pro Lys Ser Gly Pro Ser Ser Leu Ala Ile Pro Gly Ala Gln Asp Arg
305     310     315     320
Glu Glu Glu Thr Gly Gly Gly Gly Gln Asn Gly Thr Glu Val Gly Glu
325     330     335
Asp Gly Ile Gly Gln Ala Gly Asn Arg
340     345

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<210> 38
 <211> 355
 <212> PRT
 <213> Homo sapiens

<400> 38

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Met Arg Lys Gly Thr His Leu Leu Val Pro Cys Leu Glu Glu Glu Glu
1      5      10      15
Leu Ala Leu His Arg Arg Arg Leu Asp Met Ser Glu Ala Leu Pro Cys
20      25      30
Pro Gly Lys Glu Thr Pro Thr Pro Gly Cys Arg Leu Gly Ala Leu Tyr
35      40      45
Trp Ala Cys Val His Asn Asp Pro Thr Gln Leu Gln Ala Ile Leu Asp
50      55      60
Gly Gly Val Ser Pro Glu Glu Ala Thr Gln Val Asp Ser Asn Gly Arg
65      70      75      80
Thr Gly Leu Met Val Ala Cys Tyr His Gly Phe Gln Ser Val Val Ala
85      90      95
Leu Leu Ser His Cys Pro Phe Leu Asp Val Asn Gln Gln Asp Lys Gly
100     105     110
Gly Asp Thr Ala Leu Met Leu Ala Gln Ala Gly His Val Pro Leu
115     120     125
Val Ser Leu Leu Leu Asn Tyr Tyr Val Gly Leu Asp Leu Glu Arg Arg
130     135     140
Asp Gln Arg Gly Leu Thr Ala Leu Met Lys Ala Ala Met Arg Asn Arg
145     150     155     160
Cys Glu Cys Val Ala Thr Leu Leu Met Ala Gly Ala Asp Leu Thr Ala
165     170     175
Val Asp Pro Val Arg Gly Lys Thr Ala Leu Glu Trp Ala Val Leu Thr
180     185     190
Asp Ser Phe Asp Thr Val Trp Arg Ile Arg Gln Leu Leu Arg Arg Pro
195     200     205
Gln Val Glu Gln Leu Ser Gln His Tyr Lys Pro Glu Trp Pro Ala Leu
210     215     220
Ser Gly Leu Val Ala Gln Ala Gln Ala Gln Val Ala Pro Ser
225     230     235     240
Leu Leu Glu Arg Leu Gln Ala Thr Leu Ser Leu Pro Phe Ala Pro Ser
245     250     255
Pro Gln Glu Gly Gly Val Leu Asp His Leu Val Thr Ala Thr Thr Ser
260     265     270
Leu Ala Ser Pro Phe Val Thr Thr Ala Cys His Thr Leu Cys Pro Asp
275     280     285
His Pro Pro Ser Leu Gly Thr Arg Ser Lys Ser Val Pro Glu Leu Leu
290     295     300
Val Pro Ala Glu Ala Gln Ser Phe Arg Thr Pro Lys Ser Gly Pro Ser
305     310     315     320
Ser Leu Ala Ile Pro Gly Ala Gln Asp Arg Glu Glu Glu Thr Gly Gly
325     330     335
Gly Gly Gln Asn Gly Thr Glu Val Gly Glu Asp Gly Ile Gly Gln Ala
340     345     350
Gly Asn Arg
355

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<210> 39

<211> 183

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<223> genomic DNA, Exon from 1 to 143

<400> 39

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gaaagtccag ccattctgtta cctgcgttgc ttcttggggr gggatagtcc acctggaggc      60
attcggagac ccagtgtattg tgctccgygg agcctgggct gtgccccgcg ttgactgcct      120
catagatacc ctacgaaccc caagtaagaa aaaacgacga ccctctctcc gtgagtctca      180
ctg                                          183

```

<210> 40

<211> 462

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<223> genomic DNA, Exon from 108 to 358

<400> 40

```

gggataaatg ttttccctgg ggcaagggct gtgcacttgc cagctgtgtg gtccccctccc      60
taggatccag ggagacactc actactcctc tccattctgt gtttttagatg ccagctgcat      120
gagaaaaggg actcaccttc tggttccctg cctggaagag gaagagctgg cattgcacag      180
gagacggctg gacatgtctg aggcactgcc ctgcccgggc aaggagaccc ccaccccagg      240
ctgcaggctg ggggccctgt attgggcctg tgtccacaat gatccccacc agctccaagc      300
catactggat ggtggggctc ccccagagga ggccacccag gtggacagca atgggagggg      360
gagatgtcct ggcttcccag aacagctggg ggcattcttg catccccacc acaccgtcct      420
ggcctggctc cctgagaggg gttcaggggc aatacctcct gc                               462

```

<210> 41

<211> 308

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<223> genomic DNA, Exon from 89 to 218

<400> 41

```

ctctgggaca gatatgggtt tagaggggtgc aaggggccct ggagtggccc aggggggaaag      60
caggggatct gagctgcccc tccctcagac aggcctcatg gtcgcatgct accacggctt      120
ccagagtgtt gtggccctgc tcagccactg tcccttcctt gatgtgaacc agcaggacaa      180
aggaggggac acggccctca tgttggctgc ccaagcaggt gtgaggctgc tgcacccac      240
ttcogacagc ccccttttga tgcagacagg gcctcagccc cacccttggt gcacggtggt      300
ctacacca                                     308

```

<210> 42

<211> 231

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<223> genomic DNA, Exon from 49 to 159

<400> 42

```

tcatcacccc ctttcctggg gaccaagctt acccttgctg ccctgcaggc cacgtgcctc      60
tagtgagtct cctgctcaac tactatgtgg gcctggacct ggaacgccgg gaccagcggg      120

```

```

ggctcacggc gttaatgaag gctgccatgc ggaaccgctg tgagtgcgtg gccaccctcc 180
tcatggcagg tgtgcggggc ctggaccggg gtgtgtggcc tccagtcct c 231

```

```

<210> 43
<211> 231
<212> DNA
<213> Homo sapiens

```

```

<220>
<221> misc_feature
<223> genomic DNA, Exon from 49 to 189

```

```

<400> 43
tcatcacccc ctttcctggg gaccaagctt acccttgctg ccctgcaggc cacgtgcctc 60
tagtgagtct cctgctcaac tactatgtgg gcctggacct ggaacgccgg gaccagcggg 120
ggctcacggc gttaatgaag gctgccatgc ggaaccgctg tgagtgcgtg gccaccctcc 180
tcatggcagg tgtgcggggc ctggaccggg gtgtgtggcc tccagtcct c 231

```

```

<210> 44
<211> 588
<212> DNA
<213> Homo sapiens

```

```

<220>
<221> misc_feature
<223> genomic DNA, Exon from 98 to 499

```

```

<400> 44
aatgtaaccc acatcagtct tgctcctaaa gaatctgccc ttccacaaat caccaacccc 60
tatcccggcc catgtcacc cctgtgctcc ttcccagggt ctgacctgac agcagtggac 120
cctgttcggg gcaagacggc cctggaatgg gcagtgtga ccgacagctt cgacacogtg 180
tggaggattc ggcagctgct gaggcggccc caagtggagc agcttagcca gcaactacaag 240
cccagtgggc cggccttgct cgggctcgtg gcccaggccc aggccaggc ccaggttgcc 300
ccttcactcc tagaacggct gcaggctacc ttgagcctcc cctttgcccc gtctcctcag 360
gaggggggtg ttctggacca ccttgtgact gccacaacca gcctggccag tcccttcgtc 420
accactgcct gccacactct gtgccctgac catccacctt cgctgggcac ccgaagcaag 480
tccgtgccag agctgttagg tactgccccg cccctcccc tggttcccc gtecccgcca 540
gggagtcccc agaggtcccc gtgggtcttc gtcccctacc agagccct 588

```

```

<210> 45
<211> 503
<212> DNA
<213> Homo sapiens

```

```

<220>
<221> misc_feature
<223> genomic DNA, Exon from 27 to 503

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```

<400> 45
ccaaggcatc ctcatcctcc caccagtgcc agccgaagcc cagtccttca ggacaccaaa 60
gtctggccct tctctctggt cgataccagg agctcaggat agagaagagg aaacaggagg 120
aggaggccag aatggcacag aagtagggga agatgggata ggacaggctg ggaacaggta 180
atcaggcccc tcccagggtt tctttccct ctggagtgcc tccggcctcc ccatccacct 240
ctgcctaagt aaatctgctc tcaacctata tatatacaag gtcattcatt ctagcattgt 300
ttgcaagagt gaaagagtgg aaacaccga agtgtccatc agtaaggagc aggctagatt 360
gattacggat gtaattgctg tccatccata cagagcatac tctacagtgt attctaaaat 420

```

aagactaagg aagctgttta tattctgata tgaaactacc atcaagatgt ataaagtaaa 480
aataactaag gagtggaaca gtg 503

<210> 46
<211> 18
<212> DNA
<213> Artificial Sequence

<220>
<223> primer

<400> 46
ctcacatcct tctcagcc 18

<210> 47
<211> 19
<212> DNA
<213> Artificial Sequence

<220>
<223> primer

<400> 47
gtggaatgtc agggaaatc 19

<210> 48
<211> 18
<212> DNA
<213> Artificial Sequence

<220>
<223> primer

<400> 48
tgactgcctc caggaatt 18

<210> 49
<211> 18
<212> DNA
<213> Artificial Sequence

<220>
<223> primer

<400> 49
ttacgaaatg aatgggacg 18

<210> 50
<211> 18
<212> DNA
<213> Artificial Sequence

<220>

<223> primer

<400> 50

aggctctagg tccatgac

18

<210> 51

<211> 19

<212> DNA

<213> Artificial Sequence

<220>

<223> primer

<400> 51

atgtgaaatc tgcgaaagg

19

<210> 52

<211> 18

<212> DNA

<213> Artificial Sequence

<220>

<223> primer

<400> 52

cgtgccatga ctgagtac

18

<210> 53

<211> 18

<212> DNA

<213> Artificial Sequence

<220>

<223> primer

<400> 53

aactgcagtg ggtaccag

18

<210> 54

<211> 19

<212> DNA

<213> Artificial Sequence

<220>

<223> primer

<400> 54

tctgagccta gaggataacc

19

<210> 55

<211> 18

<212> DNA
 <213> Artificial Sequence

<220>
 <223> primer

<400> 55
 gatctcagag gcaggttg

18

<210> 56
 <211> 20
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> primer

<400> 56
 tgctgtgaag attggagatc

20

<210> 57
 <211> 36
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> primer

<220>
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 <222> (24)..(35)
 <223> n = inosine

<400> 57
 ggccacgcgt cgactagtac gggnnngggnn gggnnng

36

<210> 58
 <211> 20
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 <213> Artificial Sequence

<220>
 <223> primer

<400> 58
 ggccacgcgt cgactagtac

20

<210> 59
 <211> 20
 <212> DNA
 <213> Artificial Sequence

<220>

<223> primer
<400> 59
agcttgaagt ggctaaagtc 20

<210> 60
<211> 20
<212> DNA
<213> Artificial Sequence

<220>
<223> primer
<400> 60
tgatctccaa tcttcacagc 20

<210> 61
<211> 18
<212> DNA
<213> Artificial Sequence

<220>
<223> primer
<400> 61
tgtgccagga aaggaagg 18

<210> 62
<211> 19
<212> DNA
<213> Artificial Sequence

<220>
<223> primer
<400> 62
tagtcagcag catcggggg 19

<210> 63
<211> 21
<212> DNA
<213> Artificial Sequence

<220>
<223> primer
<400> 63
agcaagttca gcctgggttaa g 21

<210> 64
<211> 18
<212> DNA

<213> Artificial Sequence
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<223> primer
<400> 64
atgttcagtc aggcaggg 18

<210> 65
<211> 18
<212> DNA
<213> Artificial Sequence
<220>
<223> primer
<400> 65
ttcttgtagc atgccctc 18

<210> 66
<211> 18
<212> DNA
<213> Artificial Sequence
<220>
<223> primer
<400> 66
cggaaccgct gtgagtgc 18

<210> 67
<211> 18
<212> DNA
<213> Artificial Sequence
<220>
<223> primer
<400> 67
taggcagagg tggatggg 18

<210> 68
<211> 18
<212> DNA
<213> Artificial Sequence
<220>
<223> primer
<400> 68
ggccactcgg gctttag 18

<210> 69
<211> 18
<212> DNA
<213> Artificial Sequence

<220>
<223> primer

<400> 69
gtgcaatgcc agctcttc

18

<210> 70
<211> 18
<212> DNA
<213> Artificial Sequence

<220>
<223> primer

<400> 70
tgccaagctg ttagtgcc

18

C!
Cont
<210> 71
<211> 18
<212> DNA
<213> Artificial Sequence

<220>
<223> primer

<400> 71
catgctacca cggcttcc

18